Table 3: Chromosomal Location of DNA Probes Utilitzed

<u>Probe</u>	Locus <u>Symbol</u>	Location	Restriction <u>Enzyme</u>	Reference
рЈО71Н-А	D5S20	5p13	EcoRI	1
p105-153A	D5S39	5q11.2-q13.3	MspI	1
π227	D5S37	5q21	PstĪ	1
C11p11	D5S71	5q14-q21	TaqI	1
M4	D5S6	5q11.2-q13.3	BamHI	1
J0205H-C	D5S22	5q34-qter	MspI	1
pYNZ22	D17S30	17p13.3	BamHI	2
pYNH37.3	D17S28	17p13.3	TaqI	1
pTHH59	D17S4	17q23-q25.3	TaqI	1
L2.7	D18S6	18p11	PstĪ	1
рНН64	TTR	18q11.2-q12.1	MspI	1
DCC1.9	DCC	18q21.3	EcoRI	4
p15-65	D18S8	18q21.3	MspI	3
pERT25	D18S11	18q23	PstĬ	1

- Cytogenetics and Cell Genetics 58: 1-2200 (1991).
   Nakamura Y: Nuc Acids Res 16:4707 (1987).
- 3. Marthens F, et al. Nuc. Acids Res 15:1348 (1987).
- 4. Vogelstein, unpublished results.

WO 94/19492

Table 4: Frequency of Allelic Loss on Chromosomes 5, 17 and 18

Chromosome	Lo	oss	No Loss	NI
	No.	<b>%</b> *	(No.)	(No.)
5p	9.	20	36	46
<b>5</b> q	40	46	47	4
5	42	48	46	3
17p	59	69	26	6
17q	29	45	36	26
17	61	69	28	2
18p	24	69	11	56
18q	59	69	27	5
18	62	70	27	2

NI = non-informative

<sup>\*</sup> percentage refers to the number of tumors demonstrating loss of heterozygosity divided by the total number of tumors that were informative for the DNA probes utilized.

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Table 5: Association of DNA Alterations with Loss of Heterozygosity

		NL	L	
		5q		
MFD 27	Neg I II	34 (33) 11 (8) 1	37 (36) 1 2	p=0.025
		17p		
MFD 41	Neg I II	17 (16) 7 (6) 2	56 (55) 1 (0) 0	p<0.0005
		18q		
MFD 26	Neg I II	15 (14) 10 (8) 1	53 (51) 0 4	p<0.0005
	= no loss			

NL = no loss
L = loss
( ) synchronous tumors omitted

PCT/US94/01761

Table 6 -- Association of DNA alterations with anatomical site.

				35		
	p=0.003	p=0.018	p=0.003	p=0.006	p=0.003	_
٥	49 (47) 1 2	45 (43) 1 6	51 (50) 0 0	48 (46) 1 3	49 (48)	chronous mitted
۵.	26 11 (8) 1	25 8 (6) 5 (4)	28 (27) 8 (6) 2	25 10 (7) 2	24 13 (10)	with synchronous tumors omitted
				·		, 🗀
œ	26 (25) 1 2	23 (22) 1 5	28 0 0	27 (26) 1 1	26	
S	20 (19) 0 0	19 (18) 0 1	20 (19) 0 0	18 (17) 0 2	20 (19)	lon
OC	0 0 0	0 0 .	0 0 0	0 0 5	0 0	descending colon sigmoid rectum proximal colon distal colon
SF	0	00	-00	00	0	descen sigmoi rectum proxim
TC	5 1 (0) 0	5 1 (0) 0	6 (5) 0 0	5 1 (0) 0	5 1 (0)	OC S R G
붓	2 1 0	2 0 1	201	0		
AC	4 4 (3) 1	3 4 2 (1)	4 5 (4) 0	4 4 (3)	3 (5)	ng colon flexure rse colon flexure
ပ	15 5 (4) 0	15 3 (2) 2	16 3 (2) 1	15 4 (3) 1	15 5 (4)	caecum ascending colon hepatic flexure transverse colon splenic flexure
•	Neg I II	Neg I I I	Neg I II	Neg I II	Tumors demon- strating mutations at	AC T. T. S.
	MF027	635	MFD41	MFD26	Tumors demonstrating mutations at	

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Table 7. Clinicopathologic findings in MTS patients with tumors exhibiting microsatellite instability (Group I: cases 4, 6, 9, 23, 24, and 25) and with tumors not exhibiting microsatellite instability (Group II: cases 3, 5, 11, 17, 19, 21, and 22).

		Other skin		Visceral malignancy		
Case	Number of	tumors		v meerer ramegarana,		1
#	sebaceous	tuniois	Site	Histology	Stage	Age° 53
	tumor(s)	BCC	Cecum*	ACA	B2	53
4	5.	KA*			İ	
		SK (12)	1			
Į l		VK (14)	<u></u> i			
	7*	AK (>50)	Ascending colon	? (outside institution)	?	42
6	<b>'</b>	IFK	Rectosigmoid colon*	` ACA	B2	47
		KA	Bladder	TCC	l I	49 72
	ì	SCC (2)	Inner ear	SCC .	1	73
1		SK (3) VK	Kidney*	RCC	<b>!</b>	74
i	1	V K	Bladder	TCC RCC		74
	İ		Kidney	Metastatic ACA		74
1			Liver	ACA	В2	38
9	5*	BCC (2)	Transverse colon	CLL	]	62
1	l	KA (2)	Lymph node* Ascending colon*	ACA	B2	67
1	l .	SCC (4) SK	Lymph node	CLL		69
i		VK (3)	Lynq			
<u></u>	6*	AK (2)	Rectum*	ACA	B1	42
23	0.	scc'	Ascending colon	ACA	B2	64
		SCČ´ VK	Transverse colon	ACA	B2	65 70
		'	Kidney (renal	TCC	1	J / 0
	1	.	pelvis)*	ACA	1	73
		<u> </u>	Prostate	ACA	C2	23
24	5 *	AK (2) EC (5)	Cecum Rectum	ACA ACA	C2	29
	1	EC (5)	Liver	Metastatic ACA	l D	30
1	[	l IFK	Ovary	Adenoacanthoma	1	33
			Cecum*	ACA	B2	51
			Rectum *	ACA	B1	64
25	1*	AK (3)	Descending colon	ACA	C2	40
25	1	1	Ascending colon*	ACA	B1 D	52 62
l			Retroperitoneum	Metastatic ACA	1 D	79
3	1*	BCC (2)*	Bladder	TCC	1	'"
1		Melanoma		ì	1	1
	I	SK	Calan (anlaria	ACA	C2	69
5	1	••••	Colon (splenic flexure)	1	1	1
1	1		Colon (hepatic	ACA	C2	79
1		1	flexure)*		1	I .
1			Peritoneum	Metastatic ACA	D_	80
11	$\frac{1}{1}$	<del></del>	Cecum*	ACA	B2	79
11	1 .	''''	Lymph node (groin)	NHL		93
17	1	EC	Stomach*	NHL	i i	62
1 1	1 -	SK	Cervix	In situ SCC	1	64
	<u> </u>	<u> </u>	Stomach*	NHL		68
19	1*	EC	Breast*	ACA ACA	<del>- </del>	68
21	1	AK (2)	Ascending colon*	Metastatic ACA	A D	68
1	1	ļ	Jejunum*	Metastatic ACA Metastatic ACA	l b	68
	1		Peritoneum Breast*	Invasive ACA	<del>                                     </del>	74
22	1*		Dreast	HIVASIVETICA		

KEY TO ABBREVIATIONS AND SUMBOLS
ACA-adenocarcinoma R
AK-actinic keratosis S
BCC-basal cell carcinoma S
CLL-chronic lymphocytic leukemia T
EC-epidermoid cyst V
IFK-inverted follicular keratosis KA-keratoacanthoma NHL-non-Hodgkin's lymphoma

S
RCC-renal cell carcinoma
SCC-squamous cell carcinoma
SK-seborrheic keratosis
TCC-transitional cell carcinoma
VK-verrucal keratosis
\*PCR data obtained
†Stage of colorectal cancer
\*Age at diagnosis of visceral malignancy

Table 8. Results of assays for microsatellite instability in the various tumors.

### PATIENTS WITH MICROSATELLITE INSTABILITY +

Case #	Tumor site (histology)*	# of loci demonstrating microsatellite instability
4	Cecum (ACA) Skin (SE) Skin (KA)	4/4 4/4 4/4
6	Rectosigmoid colon (ACA) Kidney (RCC) Skin (SA) Skin (SA)	3/4 0/4 4/4 3/4
9	Skin (SA) Skin (SA) Ascending Colon (ACA) Skin (SC) Skin (SC) Lymph node (CLL)	4/4 3/4 4/4 4/4 4/4 0/4
23	Rectum (ACA) Skin (SA) Skin (SA) Renal pelvis (TCC) Skin (SC) Prostate (ACA)	4/4 4/4 4/4 3/4 4/4 3/4
24	Cecum (ACA) Skin (SA) Rectum (ACA)	3/4 4/4 3/4
25	Ascending colon (ACA) Skin (SA)	4/4 3/4

### PATIENTS WITHOUT MICROSATELLITE INSTABILITY

Case #	Tumor site (histology)	# of loci demonstrating microsatellite instability
3	Skin (BCC) SKin (SC)	0/4 0/4
5	Hepatic flexure (ACA)	0/4
11	Cecum (ACA)	0/4
17	Stomach (NHL) Stomach (NHL)	0/4 0/4
19	Eyelid (Meibomian gland adenoma) Breast (ACA)	0/4 0/4
21	Ascending colon (ACA) Jejunum (metastatic ACA)	0/4 1/4
22	Eyelid (Meibomian gland carcinoma) Breast (ACA)	0/4 0/4

KEY TO ABBREVIATIONS AND SYMBOLS
\*Tumors are ordered according to sequence of occurrence in patient.
finstability at 3 of 4 loci
SA-sebaceous adenoma
SE-sebaceous epithelioma
SC-sebaceous carcinoma
See Table 1 for abbreviations of remaining tumor types.

Table 9. Summary of clinical features of patients with or without microsatellite instability.

# PATIENTS WITH MICROSATELLITE INSTABILITY

	_	_	7	_	_	7			_		_	т-	_	7	_	7
Comments	Alive, no evidence of malignancy at age 78.	Death from renal cell carcinomia at age 74.	Death from CLL at age by.	Death from Alzheimer's disease at age 70. 140 mailghairty at ucaut.	Alive, no evidence of mangnancy at age /u.	Death from suicide at age 63. Metastatic colon cancer at death.	ТАВІГТҮ	Comments	Death from metastatic melanoma at age 89.	Death from metastatic colon cancer at age 81.	I act to follow in at 200 95 High stage fumphoma at that time.	TOTAL TO TOTAL THE BEACH OF THE STATE OF THE	Alive, no evidence of mangrancy at age co.	Death at age 79 from neart disease	Death from metastatic colon cancer at age 65.	Death from metastatic Meibomian gland carcinoma at age 73.
Family history of cancer	Yes	Yes	Yes	Yes	Yes	Yes	PATIENTS WITHOUT MICROSATELLITE INSTABILITY	Family history of cancer	Yes	Š	200	81	Š	Š	Yes	Yes
Survival following onset of first visceral malignancy (y)	24	32	32	35	37	23	PATIENTS WITHOUT	Survival following onset of first visceral malignancy (y)	6	11	11	7.5	23	11	1	-
Age of onset of first visceral malignancy (tumor type)	53 (Cecum)	42 (Ascending colon)	38 (Transverse colon)	42(Rectum)	23 (Cecum)	40 (Descending colon)		Age of onset of first visceral malignancy (tumor type)	1.44.107.00	/y (biadder)	69 (Splenic flexure)	70 (Cecum)	62 (Lymphoma)	68 (Breast)	68 (Ascending colon)	74 (Resect)
Case #	4	9	6	23	24	25		Case #	ļ	7	2	11	17	2	21	25

finsability in at least 3 of the 4 lock
Only tumor not exhibiting microsatellite instability in this patient.

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# SEQUENCE LISTING

	(1)	GENEF	RAL IN	FORMATION:					
5		(i)	APPLI	CANT: Stephen N. Thibodeau Gary D. Bren					
10		(ii)	TITLE	OF INVENTION: TUMOR-SPECIFIC GENOMIC INSTABILITY AS A PROGNOSTIC INDICATOR					
10		(iii)	(iii) NUMBER OF SEQUENCES: 18						
		(iv)	CORRE	ESPONDENCE ADDRESS:					
15			(A)	ADDRESSEE: Patterson & Keough, P.A.					
			(B)	STREET: 1200 Rand Tower 527 Marquette Avenue South					
20			(C)	CITY: Minneapolis					
			(D)	STATE: Minnesota					
25			(E)	COUNTRY: USA					
			(F)	ZIP: 55402					
		(v)	COMPU	TTER READABLE FORM:					
30			(A)	MEDIUM TYPE: Floppy disk					
			(B)	COUMPUTER: Apple Macintosh					
35			(C)	OPERATING SYSTEM: Apple Macintosh System 7.0.1					
			(D)	SOFTWARE: WordPerfect 2.1.4 for the Macintosh					
		(vi)	CURRE	INT APPLICATION DATA:					
40			(A)	APPLICATION NUMBER:					
			(B)	FILING DATE:					
45			(C)	CLASSIFICATION:					
10		(viii	.)	ATTORNEY/AGENT INFORMATION:					
			(A)	NAME: Mark S. Ellinger, Esq.					
50			(B)	REGISTRATION NUMBER: 34,812					
			(C)	REFERENCE/DOCKET NUMBER: 1144.01-WO-01					

40

# (ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 612/349-5743

5 (B) TELEFAX: 612/349-9266

	(2)	INFORMATIO	ON FOR SEQ ID NO:1
		(i) SEQU	ENCE CHARACTERISTICS:
5		(A)	LENGTH: 21
		(B)	TYPE: nucleic acid
10		(C)	STRANDEDNESS: single
10		(D)	TOPOLOGY: linear
15		(viii)	POSITION IN GENOME:
15		(A)	CHROMOSOME/SEGMENT: 5q11.2-q13.3
		(B)	MAP POSITION: D5S107
20		(xi) A SE	QUENCE DESCRIPTION: SEQ ID NO:1:
	GATC		CCAAATA C 21
25			
	(2)	INFORMATIO	ON FOR SEQ ID NO:2
		(i) SEQU	ENCE CHARACTERISTICS:
30		(A)	LENGTH: 20
		(B)	TYPE: nucleic acid
<b>3</b> 5		(C)	STRANDEDNESS: single
		(D)	TOPOLOGY: linear
40		/*************************************	POSITION IN GENOME:
40			CHROMOSOME/SEGMENT: 5q11.2-q13.3
			MAP POSITION: D5S107
45		( <i>B</i> )	MAP POSITION: D33107
		(xi) A SE	QUENCE DESCRIPTION: SEQ ID NO:2:
	GGCA'	TCAACT TGA	ACAGCAT 20

(2) INFORMATION FOR SEQ ID NO:3 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 5 (B) TYPE: nucleic acid (C) STRANDEDNESS: single 10 (D) TOPOLOGY: linear (viii) POSITION IN GENOME: 15 (A) CHROMOSOME/SEGMENT: 17p12-p11.1 (B) MAP POSITION: D17S261 20 (xi) A SEQUENCE DESCRIPTION: SEQ ID NO:3: CAGGTTCTGT CATAGGACTA 20 25 (2) INFORMATION FOR SEQ ID NO:4 (i) SEQUENCE CHARACTERISTICS: 30 (A) LENGTH: 20 (B) TYPE: nucleic acid 35 (C) STRANDEDNESS: single (D) TOPOLOGY: linear (viii) POSITION IN GENOME: 40

(xi) A SEQUENCE DESCRIPTION: SEQ ID NO:4: TTCTGGAAAC CTACTCCTGA 20

45

(B) MAP POSITION: D17S261

(A) CHROMOSOME/SEGMENT: 17p12-p11.1

	(2)	INFORMATI	ON FOR SEQ ID NO:5
		(i) SEQ	JENCE CHARACTERISTICS:
5		(A)	LENGTH: 20
		(B)	TYPE: nucleic acid
10		(C)	STRANDEDNESS: single
10		(D)	TOPOLOGY: linear
15		(viii)	POSITION IN GENOME:
15		(A)	CHROMOSOME/SEGMENT: 18q
		(B)	MAP POSITION: D18S34
20		(xi) A SI	EQUENCE DESCRIPTION: SEQ ID NO:5:
	CAGA	AAATTC TCI	CTGGCTA 20
25			
	(2)	INFORMAT	ON FOR SEQ ID NO:6
00		(i) SEQ	JENCE CHARACTERISTICS:
30		(A)	LENGTH: 20
		(B)	TYPE: nucleic acid
35		(C)	STRANDEDNESS: single
		(D)	TOPOLOGY: linear
<b>4</b> 0		(viii)	POSITION IN GENOME:
		(A)	CHROMOSOME/SEGMENT: 18q
45		(B)	MAP POSITION: D18S34
		(xi) A S	EQUENCE DESCRIPTION: SEQ ID NO:6:
	CTCA	መረጥመረር ጥርረ	2CAAGAAT 20

	(2)	44 INFORMATION FOR SEQ ID NO:7
		(i) SEQUENCE CHARACTERISTICS:
5		(A) LENGTH: 20
		(B) TYPE: nucleic acid
10	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
		(viii) POSITION IN GENOME:
15	(A) CHROMOSOME/SEGMENT: 15q11-qter	

- (xi) A SEQUENCE DESCRIPTION: SEQ ID NO:7:
- TTGACCTGAA TGCACTGTCA 20
- 25 (2) INFORMATION FOR SEQ ID NO:8
  - (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 20
- 30 (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
- 35 (D) TOPOLOGY: linear
  - (viii) POSITION IN GENOME:
- 40 (A) CHROMOSOME/SEGMENT: 15q11-qter
- (xi) A SEQUENCE DESCRIPTION: SEQ ID NO:8:
- 45 TTCCATACCT GGCAACGAGT 20

	(2)	INFO	RMATI	ON FOR SEQ ID NO:9
		(i)	SEQU	ENCE CHARACTERISTICS:
5			(A) (B)	LENGTH: 20 TYPE: nucleic acid
			(C)	STRANDEDNESS: single
10			(D)	TOPOLOGY: linear
		(x)	PUBL	ICATION INFORMATION:
15			(A)	AUTHORS: Peterson, M.G. Tanese, N. Pugh, B.F. Tjian, R.
20			(B)	TITLE: Functional domains and upstream activation properties of cloned human TATA binding protein
25			(C)	JOURNAL: Science
23			(D)	VOLUME: 248
			(F)	PAGES: 1625-1630
30			(G)	DATE: 1990
			(K)	RELEVANT RESIDUES: 352 - 371
35		(xi)	A SE	QUENCE DESCRIPTION: SEQ ID NO:9:
	ACTG	ACCCC <i>I</i>	A CAG	CCTATTC 20
40	(2)	INFO	RMATI	ON FOR SEQ ID NO:10
		(i)	SEQU	ENCE CHARACTERISTICS:
45			(A)	LENGTH: 20
			(B)	TYPE: nucleic acid
50			(C)	STRANDEDNESS: single
			(D)	TOPOLOGY: linear

	(x)	PUBL	ICATION INFORMATION:
5		(A)	AUTHORS: Peterson, M.G. Tanese, N. Pugh, B.F. Tjian, R.
10		(B)	TITLE: Functional domains and upstream activation properties of cloned human TATA binding protein
		(C)	JOURNAL: Science
		(D)	VOLUME: 248
15		(F)	PAGES: 1625-1630
		(G)	DATE: 1990
20		(K)	RELEVANT RESIDUES: 618 - 637
25	(xi)		QUENCE DESCRIPTION: SEQ ID NO:10:
	(2) INFO	RMATI	ON FOR SEQ ID NO:11
30	(i)	SEQU	ENCE CHARACTERISTICS:
		(A)	LENGTH: 21
35		(B)	TYPE: nucleic acid
		(C)	STRANDEDNESS: single
<del>1</del> 0		(D)	TOPOLOGY: linear
	(vii:	i)	POSITION IN GENOME:
		(A)	CHROMOSOME/SEGMENT: 5q
45	(x)	PUBL	ICATION INFORMATION:
		(A)	AUTHORS: Spiro, L. et al.
50		(B)	TITLE: A CA repeat 30-70 Kb downstream from the adenomatous polyposis coli (APC) gene
		(C)	JOURNAL: Nucleic Acids Res.
55		(D)	VOLUME: 19

47

(F) PAGES: 6348 et. seq.

(G) DATE: 1991

(xi) A SEQUENCE DESCRIPTION: SEQ ID NO:11:

ACTCACTCTA GTGATAAATC G 21

	(2)	) INFORMATION FOR SEQ ID NO:12				
		(i)	SEQU	ENCE CHARACTERISTICS:		
5			(A)	LENGTH: 25		
			(B)	TYPE: nucleic acid		
4.0			(C)	STRANDEDNESS: single		
10			(D)	TOPOLOGY: linear		
15		(vii:	i)	POSITION IN GENOME:		
			(A)	CHROMOSOME/SEGMENT: 5q		
		(x)	PUBL	ICATION INFORMATION:		
20			(A)	AUTHORS: Spiro, L. et al.		
			(B)	TITLE: A CA repeat 30-70 Kb downstream from the adenomatous polyposis coli (APC) gene		
25			(C)	JOURNAL: Nucleic Acids Res.		
			(D)	VOLUME: 19		
20			(F)	PAGES: 6348 et. seq.		
30			(G)	DATE: 1991		
		(xi)	A SE	QUENCE DESCRIPTION: SEQ ID NO:12:		
35	AGCA	AGCAGATAAG ACAGTATTAC TAGTT 25				
40	(2)			ON FOR SEQ ID NO:13		
		(i)	SEQU	ENCE CHARACTERISTICS:		
<b>4</b> 5			(A)	LENGTH: 20		
			(B)	TYPE: nucleic acid		
			(C)	STRANDEDNESS: single		
50			(D)	TOPOLOGY: linear		
		(vii	i)	POSITION IN GENOME:		
55			(A)	CHROMOSOME/SEGMENT: 15q		

49

PUBLICATION INFORMATION: (x)

AUTHORS: Thibodeau, S.N. et al. (A)

TITLE: Microsatellite instability in cancer of the proximal colon (B)

JOURNAL: Science (C) 10

(D) VOLUME: 260

5

(F) PAGES: 816-819

15 (G) DATE: 1993

(xi) A SEQUENCE DESCRIPTION: SEQ ID NO:13:

20 TTGACCTGAA TGCACTGTGA 20

	(2)	INFO	RMATI	ON FOR SEQ ID NO:14	
		(i)	SEQU	ENCE CHARACTERISTICS:	
5			(A)	LENGTH: 20	
			(B)	TYPE: nucleic acid	
10			(C)	STRANDEDNESS: single	
			(D)	TOPOLOGY: linear	
		(viii)		POSITION IN GENOME:	
15			(A)	CHROMOSOME/SEGMENT: 15q	
		(x)	PUBL	ICATION INFORMATION:	
20			(A)	AUTHORS: Thibodeau, S.N. et al.	
			(B)	TITLE: Microsatellite instability in cancer of the proximal colon	
25			(C)	JOURNAL: Science	
			(D)	VOLUME: 260	
20			(F)	PAGES: 816-819	
30			(G)	DATE: 1993	
0.5		(xi)	A SE	QUENCE DESCRIPTION: SEQ ID NO:14:	
35	TTCC.	TCCATACCT GGGAACGAGT 20			
<b>4</b> 0	(2)	INFO	RMATI	ON FOR SEQ ID NO:15	
		(i)	SEQU	ENCE CHARACTERISTICS:	
			(A)	LENGTH: 24	
<b>4</b> 5			(B)	TYPE: nucleic acid	
			(C)	STRANDEDNESS: single	
50			(D)	TOPOLOGY: linear	
		(vii	i)	POSITION IN GENOME:	
55			(A)	CHROMOSOME/SEGMENT: 17p	

51

## (x) PUBLICATION INFORMATION:

- (A) AUTHORS: Jones, M.H., and Nakamura, Y.
- 5 (B) TITLE: Detection of loss of heterozygosity at the human *TP53* locus using a dinucleotide repeat polymorphism
  - (C) JOURNAL: Genes Chrom. Cancer

10 (D) VOLUME: 5

(F) PAGES: 89-90

15 (G) DATE: 1992

(xi) A SEQUENCE DESCRIPTION: SEQ ID NO:15:

AGGGATACTA TTCAGCCCGA GGTG 24